

XM048113	1	-----
ANH401	1	MAAVSLRLGDLVWGKLGRIYPPWPGKIVNPPKDLKKPRGKKCFVVKFFGTEDHAWIKVEQL
AF326966	1	MAAVSLRLGDLVWGKLGRIYPPWPGKIVNPPKDLKKPRGKKCFVVKFFGTEDHAWIKVEQL
XM048113	1	-----
ANH401	61	KPYHAHKEEMIKINKGKRFQQAQVDAVEEFLRRAGKDKQTSSHNSSDDKNRRNSSEERSRP
AF326966	61	KPYHAHKEEMIKINKGKRFQQAQVDAVEEFLRRAGKDKQTSSHNSSDDKNRRNSSEERSRP
XM048113	1	-----
ANH401	121	NSGDEKRKLSLSEGKVKKNMGEKGRVSSGSSSERGSKSPLKRAQEQQSPRKRGRPPKDEKD
AF326966	121	NSGDEKRKLSLSEGKVKKNMGEKGRVSSGSSSERGSKSPLKRAQEQQSPRKRGRPPKDEKD
XM048113	1	-----
ANH401	181	LTIPESSTVKGMMAGPMAAFKWQPTASEPVKDADPHFHHFLLSQTEKPAVCYQAITKKLK
AF326966	181	LTIPESSTVKGMMAGPMAAFKWQPTASEPVKDADPHFHHFLLSQTEKPAVCYQAITKKLK
XM048113	1	-----MGSGIVSNLLKMGHTVTVWNRTA
ANH401	241	ICEEETGSTSIQAADSTAVNGSITPTDKKIGFLGLGLMGSGIVSNLLKMGHTVTVWNRTA
AF326966	241	ICEEETGSTSIQAADSTAVNGSITPTDKKIGFLGLGLMGSGIVSNLLKMGHTVTVWNRTA
XM048113	24	EKCDLFIQEGARLGRTPAEVVSTCDITFACVSDPKAAKDLVLGPGSVLQGIRPGKCYVDM
ANH401	301	EKCDLFIQEGARLGRTPAEVVSTCDITFACVSDPKAAKDLVLGPGSVLQGIRPGKCYVDM
AF326966	301	EK-----EGARLGRTPAEVVSTCDITFACVSDPKAAKDLVLGPGSVLQGIRPGKCYVDM
XM048113	84	STVDADTVTELAQVIVSRGGRFLEAPVSGNQQLSNDGMLVILAAGDRGLYEDCSSCFQAM
ANH401	361	STVDADTVTELAQVIVSRGGRFLEAPVSGNQQLSNDGMLVILAAGDRGLYEDCSSCFQAM
AF326966	355	STVDADTVTELAQVIVSRGGRFLEAPVSGNQQLSNDGMLVILAAGDRGLYEDCSSCFQAM
XM048113	144	GKTSFFLGEVGNAAKMMLIVNMVQGSFMATIAEGLTLAHVTGQSQQTLDDILNQGLASI
ANH401	421	GKTSFFLGEVGNAAKMMLIVNMVQGSFMATIAEGLTLAHVTGQSQQTLDDILNQGLASI
AF326966	415	GKTSFFLGEVGNAAKMMLIVNMVQGSFMATIAEGLTLAQVTGQSQQTLDDILNQGLASI
XM048113	204	FLDQKQCQNILQGNFKPDFYLYKIQKDLRLAIALGDAVNHPTPMAAAANEVYKRAKALDQS
ANH401	481	FLDQKQCQNILQGNFKPDFYLYKIQKDLRLAIALGDAVNHPTPMAAAANEVYKRAKALDQS
AF326966	475	FLDQKQCQNILQGNFKPDFYLYKIQKDLRLAIALGDAVNHPTPMAAAANEVYKRAKALDQS
XM048113	264	DNDMSAVYRAYIH
ANH401	541	DNDMSAVYRAYIH
AF326966	535	DNDMSAVYRAYIH

Fig. 1